

Sequence Listing

<110> Botstein,David

Desnoyers,Luc

Ferrara,Napoleone

Fong,Sherman

Gao,Wei-Qiang

Goddard,Audrey

Gurney,Austin L.

Pan,James

Roy,Margaret Ann

Stewart,Timothy A.

Tumas,Daniel

Watanabe,Colin K.

Wood,William I.

<120> Secreted and Transmembrane Polypeptides and Nucleic
Acids Encoding the Same

<130> P2930R1C2

<150> 60/095,325

<151> 1998-08-04

<150> 60/112,851

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<151> 2000-02-11

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35 40 45
Phe Ala Ile Ala Arg Arg Leu Ala Gln Asp Gly Ala His Val Val
50 55 60
Val Ser Ser Arg Lys Gln Gln Asn Val Asp Gln Ala Val Ala Thr
65 70 75

Leu Gln Gly Glu Gly Leu Ser Val Thr Gly Thr Val Cys His Val
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 Gly Lys Ala Glu Asp Arg Glu Arg Leu Val Ala Thr Ala Val Lys
 95 100 105
 Leu His Gly Gly Ile Asp Ile Leu Val Ser Asn Ala Ala Val Asn
 110 115 120
 Pro Phe Phe Gly Ser Ile Met Asp Val Thr Glu Glu Val Trp Asp
 125 130 135
 Lys Thr Leu Asp Ile Asn Val Lys Ala Pro Ala Leu Met Thr Lys
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 Ala Val Val Pro Glu Met Glu Lys Arg Gly Gly Gly Ser Val Val
 155 160 165
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 200 205 210
 Ala Pro Gly Leu Ile Lys Thr Ser Phe Ser Arg Met Leu Trp Met
 215 220 225
 Asp Lys Glu Lys Glu Glu Ser Met Lys Glu Thr Leu Arg Ile Arg
 230 235 240
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Glu	His	Leu	Glu	Lys 470	Glu	Leu	Ser	Glu	Lys 475	Ser	Gly	Gln	Leu	Arg 480
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Ile	Asp	Gly	Lys	Leu 515	Arg	Gln	Gly	Ser	Leu 520	Leu	Ser	Pro	Glu	Glu 525
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Ala	Ala	Ile	Glu	Tyr 545	Lys	Asn	Glu	Ala	Ile 550	Thr	Cys	Arg	Gln	Arg 555
Val	Leu	Arg	Ala	Ser 560	Ala	Ser	Leu	Leu	Ser 565	Gln	Cys	Glu	Met	Asn 570
Leu	Met	Ala	Lys	Leu 575	Ser	Tyr	Leu	Ser	Ser 580	Ser	Glu	Thr	Arg	Ala 585
Leu	Leu	Cys	Lys	Tyr 590	Phe	Asp	Lys	Val	Val 595	Thr	Leu	Arg	Glu	Glu 600
Gln	His	Gln	Gln	Gln 605	Ile	Ala	Phe	Ser	Glu 610	Leu	Glu	Met	Gln	Leu 615
Glu	Glu	Gln	Gln	Arg 620	Leu	Val	Tyr	Trp	Leu 625	Glu	Val	Ala	Leu	Glu 630
Arg	Gln	Arg	Leu	Glu 635	Met	Asp	Arg	Gln	Leu 640	Thr	Leu	Gln	Gln	Lys 645
Glu	His	Glu	Gln	Asn 650	Met	Gln	Leu	Leu	Leu 655	Gln	Gln	Ser	Arg	Asp 660
His	Leu	Gly	Glu	Gly 665	Leu	Ala	Asp	Ser	Arg 670	Arg	Gln	Tyr	Glu	Ala 675
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35 40 45
Lys Ile Tyr Asn Pro Ser Glu Gln Cys Cys Tyr Asp Asp Ala Ile
50 55 60
Leu Ser Leu Lys Glu Thr Arg Arg Cys Gly Ser Thr Cys Thr Phe
65 70 75
Trp Pro Cys Phe Glu Leu Cys Cys Pro Glu Ser Phe Gly Pro Gln
80 85 90
Gln Lys Phe Leu Val Lys Leu Arg Val Leu Gly Met Lys Ser Gln
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ggcgccaggg tggcccccg ggcgcgcttg gtctcggaga agcggggacg 200
aggccggagg atgagcgact gagggcgacg cgggcactga cgcgagttgg 250
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ccctagcccc gtgtgcgcgc caggcggagc gcgcaggtgg ggctgggctg 350

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Gln Thr Glu Cys Val Asp Pro Val Tyr Glu Pro Asp Gln Cys Cys		
260	265	270
Pro Ile Cys Lys Asn Gly Pro Asn Cys Phe Ala Glu Thr Ala Val		
275	280	285
Ile Pro Ala Gly Arg Glu Val Lys Thr Asp Glu Cys Thr Ile Cys		
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Cys Thr Glu Ala Lys Asp Gln Glu Pro Arg Val Thr Glu His Arg
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Gln Glu Asp Phe Cys Asn Asn Leu Val Asn Ser Leu Pro Leu Trp
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Cys Pro Lys Gly Thr Thr His Cys Tyr Asp Gly Leu Leu Arg Leu
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Arg Gly Gly Gly Ile Phe Ser Asn Leu Arg Val Gln Gly Cys Met
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Pro Gln Pro Gly Cys Asn Leu Leu Asn Gly Thr Gln Glu Ile Gly
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Pro Val Gly Met Thr Glu Asn Cys Asn Arg Lys Asp Phe Leu Thr
200 205 210
Cys His Arg Gly Thr Thr Ile Met Thr His Gly Asn Leu Ala Gln
215 220 225
Glu Pro Thr Asp Trp Thr Thr Ser Asn Thr Glu Met Cys Glu Val

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Pro Ala Leu Phe Ser Arg Glu Asn Tyr	Gly Arg Leu Arg Leu Ile	
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Thr Ser Ser Lys His Arg Cys Met Asp	Ser Ser Ala Ala Phe Leu	
185	190	195
Gln Gly Leu Trp Gln His Tyr His Pro	Gly Leu Pro Pro Pro Asp	
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Val Ala Asp Met Glu Phe Gly Pro Pro	Thr Val Asn Asp Lys Leu	
215	220	225
Met Arg Phe Phe Asp His Cys Glu Lys	Phe Leu Thr Glu Val Glu	
230	235	240
Lys Asn Ala Thr Ala Leu Tyr His Val	Glu Ala Phe Lys Thr Gly	
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Pro Glu Met Gln Asn Ile Leu Lys Lys	Val Ala Ala Thr Leu Gln	
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Val Pro Val Asn Asp Leu Asn Ala Asp	Leu Ile Gln Val Ala Phe	
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Phe Thr Cys Ser Phe Asp Leu Ala Ile	Lys Gly Val Lys Ser Pro	
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Trp Cys Asp Val Phe Asp Ile Asp Asp	Ala Lys Val Leu Glu Tyr	
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Leu Asn Asp Leu Lys Gln Tyr Trp Lys	Arg Gly Tyr Gly Tyr Thr	
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Ile Asn Ser Arg Ser Ser Cys Thr Leu	Phe Gln Asp Ile Phe Gln	
335	340	345
His Leu Asp Lys Ala Val Glu Gln Lys	Gln Arg Ser Gln Pro Ile	
350	355	360
Ser Ser Pro Val Ile Leu Gln Phe Gly	His Ala Glu Thr Leu Leu	
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Pro Leu Leu Ser Leu Met Gly Tyr Phe	Lys Asp Lys Glu Pro Leu	
380	385	390
Thr Ala Tyr Asn Tyr Lys Lys Gln Met	His Arg Lys Phe Arg Ser	
395	400	405

Gly	Leu	Ile	Val	Pro	Tyr	Ala	Ser	Asn	Leu	Ile	Phe	Val	Leu	Tyr	410	415	420
His	Cys	Glu	Asn	Ala	Lys	Thr	Pro	Lys	Glu	Gln	Phe	Arg	Val	Gln	425	430	435
Met	Leu	Leu	Asn	Glu	Lys	Val	Leu	Pro	Leu	Ala	Tyr	Ser	Gln	Glu	440	445	450
Thr	Val	Ser	Phe	Tyr	Glu	Asp	Leu	Lys	Asn	His	Tyr	Lys	Asp	Ile	455	460	465
Leu	Gln	Ser	Cys	Gln	Thr	Ser	Glu	Glu	Cys	Glu	Leu	Ala	Arg	Ala	470	475	480
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295

300

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 305 310

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 <211> 3437
 <212> DNA
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 <211> 1029
 <212> PRT
 <213> Homo sapiens

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 35 40 45
 Asp Thr Ser Glu Glu Ala Leu Leu Leu Pro Asp Trp Leu Lys Leu
 50 55 60
 Arg Met Ile Arg Ser Glu Val Leu Arg Leu Val Asp Ala Ala Leu
 65 70 75
 Gln Asp Leu Glu Pro Gln Gln Leu Leu Leu Phe Val Gln Ser Phe
 80 85 90
 Gly Ile Pro Val Ser Ser Met Ser Lys Leu Leu Gln Phe Leu Asp
 95 100 105

				395					400					405
Thr	Ala	Asp	Ala	Ala 410	Ser	Pro	Phe	Pro	Ala 415	Cys	Lys	Pro	Val	Val 420
Val	Val	Ser	Ser	Leu 425	Leu	Leu	Gln	Glu	Glu 430	Glu	Pro	Leu	Ala	Gly 435
Gly	Lys	Pro	Gly	Ala 440	Asp	Gly	Gly	Ser	Leu 445	Glu	Ala	Val	Arg	Leu 450
Gly	Pro	Ser	Ser	Gly 455	Leu	Leu	Val	Asp	Trp 460	Leu	Glu	Met	Leu	Asp 465
Pro	Glu	Val	Val	Ser 470	Ser	Cys	Pro	Asp	Leu 475	Gln	Leu	Arg	Leu	Leu 480
Phe	Ser	Arg	Arg	Lys 485	Gly	Lys	Gly	Gln	Ala 490	Gln	Val	Pro	Ser	Phe 495
Arg	Pro	Tyr	Leu	Leu 500	Thr	Leu	Phe	Thr	His 505	Gln	Ser	Ser	Trp	Pro 510
Thr	Leu	His	Gln	Cys 515	Ile	Arg	Val	Leu	Leu 520	Gly	Lys	Ser	Arg	Glu 525
Gln	Arg	Phe	Asp	Pro 530	Ser	Ala	Ser	Leu	Asp 535	Phe	Leu	Trp	Ala	Cys 540
Ile	His	Val	Pro	Arg 545	Ile	Trp	Gln	Gly	Arg 550	Asp	Gln	Arg	Thr	Pro 555
Gln	Lys	Arg	Arg	Glu 560	Glu	Leu	Val	Leu	Arg 565	Val	Gln	Gly	Pro	Glu 570
Leu	Ile	Ser	Leu	Val 575	Glu	Leu	Ile	Leu	Ala 580	Glu	Ala	Glu	Thr	Arg 585
Ser	Gln	Asp	Gly	Asp 590	Thr	Ala	Ala	Cys	Ser 595	Leu	Ile	Gln	Ala	Arg 600
Leu	Pro	Leu	Leu	Leu 605	Ser	Cys	Cys	Cys	Gly 610	Asp	Asp	Glu	Ser	Val 615
Arg	Lys	Val	Thr	Glu 620	His	Leu	Ser	Gly	Cys 625	Ile	Gln	Gln	Trp	Gly 630
Asp	Ser	Val	Leu	Gly 635	Arg	Arg	Cys	Arg	Asp 640	Leu	Leu	Leu	Gln	Leu 645
Tyr	Leu	Gln	Arg	Pro 650	Glu	Leu	Arg	Val	Pro 655	Val	Pro	Glu	Val	Leu 660
Leu	His	Ser	Glu	Gly 665	Ala	Ala	Ser	Ser	Ser 670	Val	Cys	Lys	Leu	Asp 675
Gly	Leu	Ile	His	Arg 680	Phe	Ile	Thr	Leu	Leu 685	Ala	Asp	Thr	Ser	Asp 690

Ser Arg Ala Leu Glu Asn Arg Gly Ala Asp Ala Ser Met Ala Cys	695	700	705
Arg Lys Leu Ala Val Ala His Pro Leu Leu Leu Arg His Leu	710	715	720
Pro Met Ile Ala Ala Leu Leu His Gly Arg Thr His Leu Asn Phe	725	730	735
Gln Glu Phe Arg Gln Gln Asn His Leu Ser Cys Phe Leu His Val	740	745	750
Leu Gly Leu Leu Glu Leu Leu Gln Pro His Val Phe Arg Ser Glu	755	760	765
His Gln Gly Ala Leu Trp Asp Cys Leu Leu Ser Phe Ile Arg Leu	770	775	780
Leu Leu Asn Tyr Arg Lys Ser Ser Arg His Leu Ala Ala Phe Ile	785	790	795
Asn Lys Phe Val Gln Phe Ile His Lys Tyr Ile Thr Tyr Asn Ala	800	805	810
Pro Ala Ala Ile Ser Phe Leu Gln Lys His Ala Asp Pro Leu His	815	820	825
Asp Leu Ser Phe Asp Asn Ser Asp Leu Val Met Leu Lys Ser Leu	830	835	840
Leu Ala Gly Leu Ser Leu Pro Ser Arg Asp Asp Arg Thr Asp Arg	845	850	855
Gly Leu Asp Glu Glu Gly Glu Glu Glu Ser Ser Ala Gly Ser Leu	860	865	870
Pro Leu Val Ser Val Ser Leu Phe Thr Pro Leu Thr Ala Ala Glu	875	880	885
Met Ala Pro Tyr Met Lys Arg Leu Ser Arg Gly Gln Thr Val Glu	890	895	900
Asp Leu Leu Glu Val Leu Ser Asp Ile Asp Glu Met Ser Arg Arg	905	910	915
Arg Pro Glu Ile Leu Ser Phe Phe Ser Thr Asn Leu Gln Arg Leu	920	925	930
Met Ser Ser Ala Glu Glu Cys Cys Arg Asn Leu Ala Phe Ser Leu	935	940	945
Ala Leu Arg Ser Met Gln Asn Ser Pro Ser Ile Ala Ala Ala Phe	950	955	960
Leu Pro Thr Phe Met Tyr Cys Leu Gly Ser Gln Asp Phe Glu Val	965	970	975
Val Gln Thr Ala Leu Arg Asn Leu Pro Glu Tyr Ala Leu Leu Cys			

980

985

990

Gln Glu His Ala Ala Val Leu Leu His Arg Ala Phe Leu Val Gly
995 1000 1005

Met Tyr Gly Gln Met Asp Pro Ser Ala Gln Ile Ser Glu Ala Leu
1010 1015 1020

Arg Ile Leu His Met Glu Ala Val Met
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<211> 2186

<212> DNA

<213> Homo sapiens

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acgttggtgga cgtttaccag cgggagttcc tggcgctgcg cgatcggttg 200
cacgcagctg agcaggagag cctcaagcgc tccaaggagc tcaacctggt 250
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<211> 548
<212> PRT
<213> Homo sapiens

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20 25 30

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Leu	Arg	Asp	Arg	Leu	His	Ala	Ala	Glu	Gln	Glu	Ser	Leu	Lys	Arg		50	55	60
Ser	Lys	Glu	Leu	Asn	Leu	Val	Leu	Asp	Glu	Ile	Lys	Arg	Ala	Val		65	70	75
Ser	Glu	Arg	Gln	Ala	Leu	Arg	Asp	Gly	Asp	Gly	Asn	Arg	Thr	Trp		80	85	90
Gly	Arg	Leu	Thr	Glu	Asp	Pro	Arg	Leu	Lys	Pro	Trp	Asn	Gly	Ser		95	100	105
His	Arg	His	Val	Leu	His	Leu	Pro	Thr	Val	Phe	His	His	Leu	Pro		110	115	120
His	Leu	Leu	Ala	Lys	Glu	Ser	Ser	Leu	Gln	Pro	Ala	Val	Arg	Val		125	130	135
Gly	Gln	Gly	Arg	Thr	Gly	Val	Ser	Val	Val	Met	Gly	Ile	Pro	Ser		140	145	150
Val	Arg	Arg	Glu	Val	His	Ser	Tyr	Leu	Thr	Asp	Thr	Leu	His	Ser		155	160	165
Leu	Ile	Ser	Glu	Leu	Ser	Pro	Gln	Glu	Lys	Glu	Asp	Ser	Val	Ile		170	175	180
Val	Val	Leu	Ile	Ala	Glu	Thr	Asp	Ser	Gln	Tyr	Thr	Ser	Ala	Val		185	190	195
Thr	Glu	Asn	Ile	Lys	Ala	Leu	Phe	Pro	Thr	Glu	Ile	His	Ser	Gly		200	205	210
Leu	Leu	Glu	Val	Ile	Ser	Pro	Ser	Pro	His	Phe	Tyr	Pro	Asp	Phe		215	220	225
Ser	Arg	Leu	Arg	Glu	Ser	Phe	Gly	Asp	Pro	Lys	Glu	Arg	Val	Arg		230	235	240
Trp	Arg	Thr	Lys	Gln	Asn	Leu	Asp	Tyr	Cys	Phe	Leu	Met	Met	Tyr		245	250	255
Ala	Gln	Ser	Lys	Gly	Ile	Tyr	Tyr	Val	Gln	Leu	Glu	Asp	Asp	Ile		260	265	270
Val	Ala	Lys	Pro	Asn	Tyr	Leu	Ser	Thr	Met	Lys	Asn	Phe	Ala	Leu		275	280	285
Gln	Gln	Pro	Ser	Glu	Asp	Trp	Met	Ile	Leu	Glu	Phe	Ser	Gln	Leu		290	295	300
Gly	Phe	Ile	Gly	Lys	Met	Phe	Lys	Ser	Leu	Asp	Leu	Ser	Leu	Ile		305	310	315
Val	Glu	Phe	Ile	Leu	Met	Phe	Tyr	Arg	Asp	Lys	Pro	Ile	Asp	Trp				

<210> 26
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 <213> Artificial Sequence

 <220>
 <223> Synthetic Oligonucleotide Probe

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 <210> 27
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 <220>
 <223> Synthetic Oligonucleotide Probe

 <400> 27
 actcgggatt cctgctgtt 19

 <210> 28
 <211> 23
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 <220>
 <223> Synthetic Oligonucleotide Probe

 <400> 28
 aggcctttac ccaaggccac aac 23

 <210> 29
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 <400> 29
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 <210> 30
 <211> 22
 <212> DNA
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 <223> Synthetic Oligonucleotide Probe

 <400> 30
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 <212> DNA

<213> Artificial Sequence

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<223> Synthetic Oligonucleotide Probe

<400> 31

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<210> 32

<211> 23

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<220>

<223> Synthetic Oligonucleotide Probe

<400> 32

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<210> 33

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 33

gtccagcaag ccctcatt 18

<210> 34

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 34

cttctgggcc acagccctgc 20

<210> 35

<211> 21

<212> DNA

<213> Artificial Sequence

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<223> Synthetic Oligonucleotide Probe

<400> 35

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<223> Synthetic Oligonucleotide Probe

<400> 36

ccagtcaggc cgttttaga 19

<210> 37

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 37

cgggcgcca agtaaaagct c 21

<210> 38

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic Oligonucleotide Probe

<400> 38

cataaagtag tatatgcatt ccagtgtt 28

TOGETHER